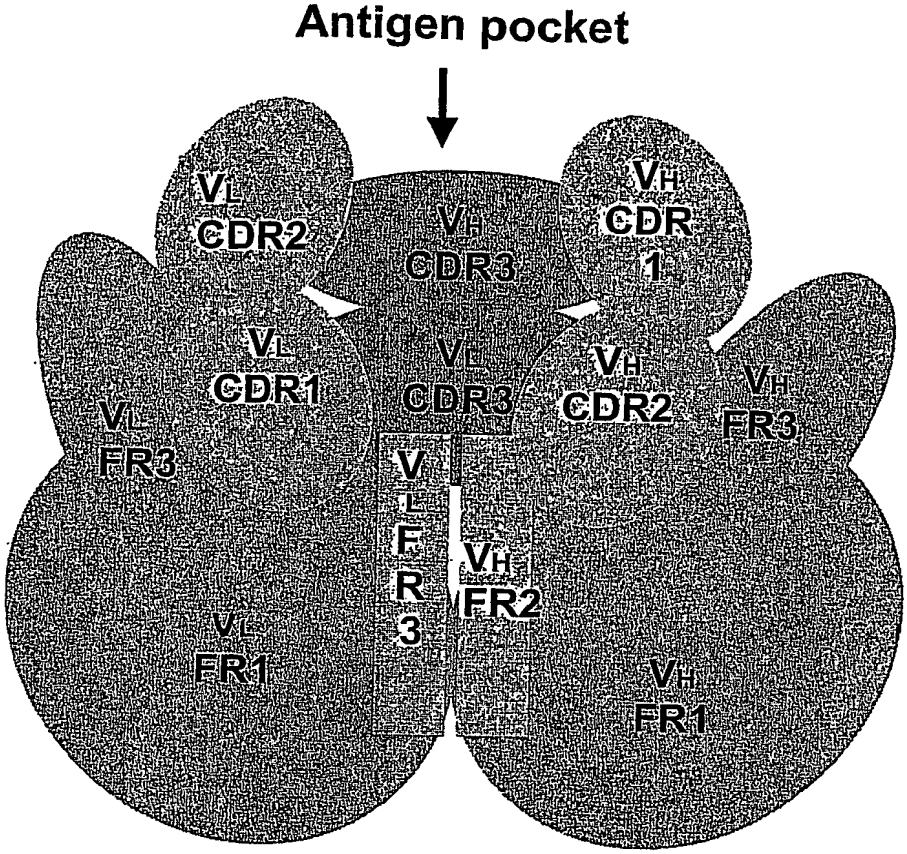
Fig. 1

Fa



CDR: complementarity determining region
FR: frame region
V_L: variabel light chain
V_H: variabel heavy chain

Fig. 2a

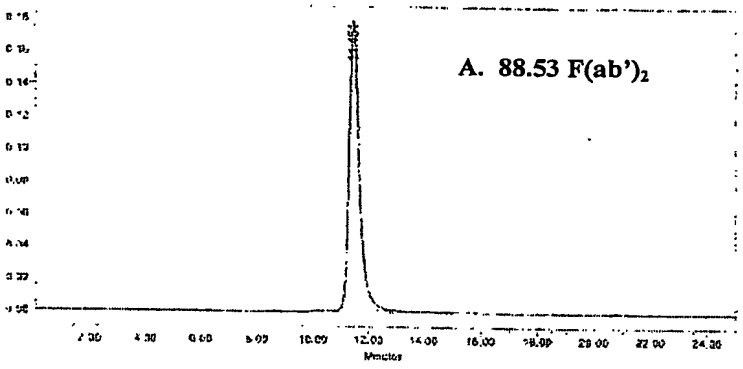


Fig. 2b

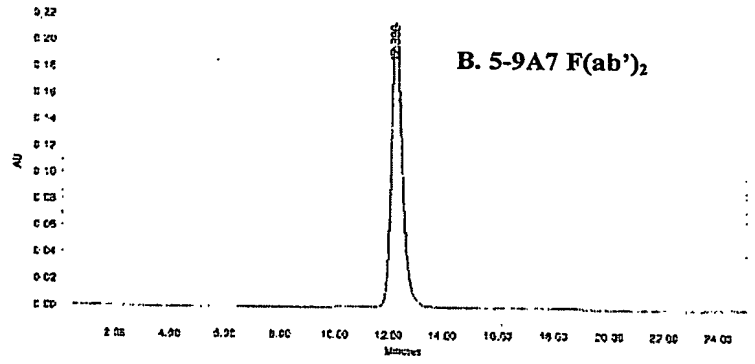


Fig. 3

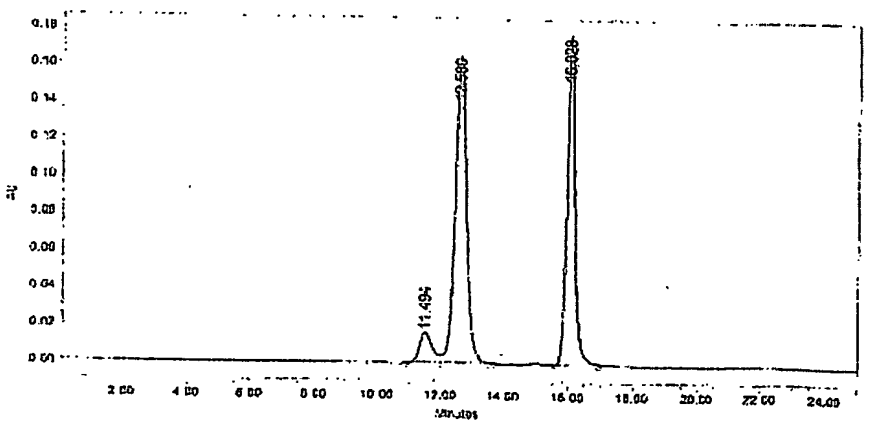


Fig. 4

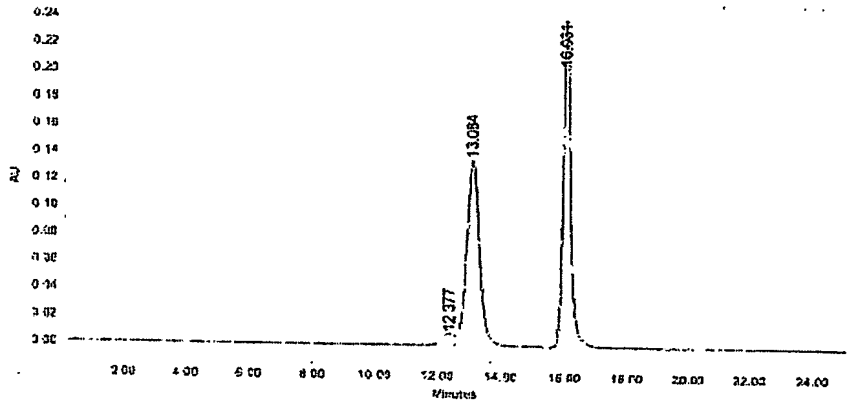


Fig. 5

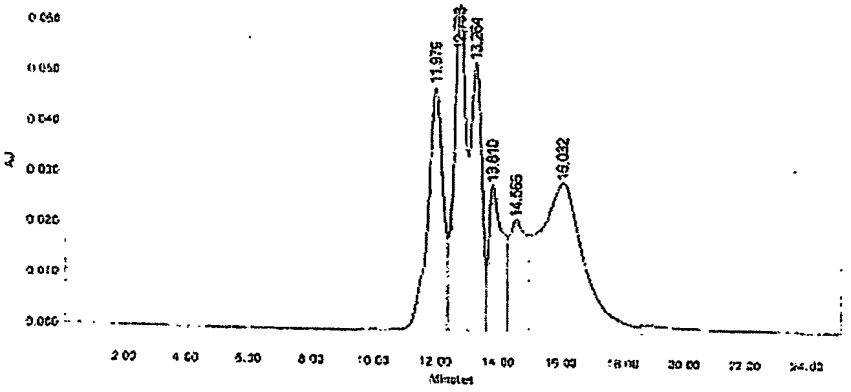


Fig. 6

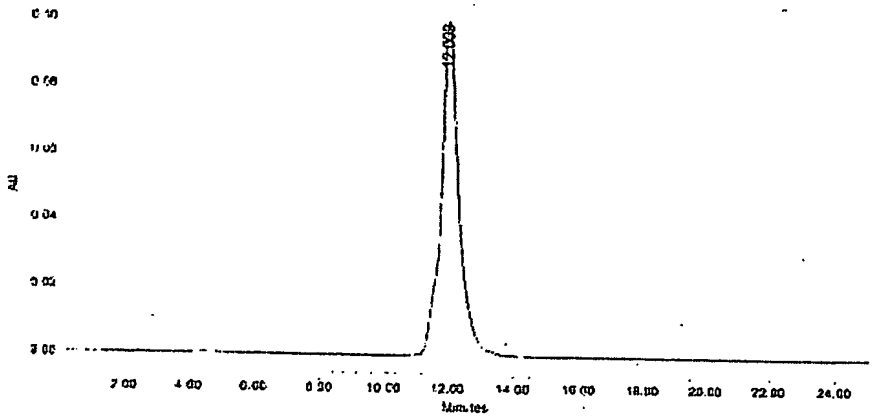


Fig. 7a

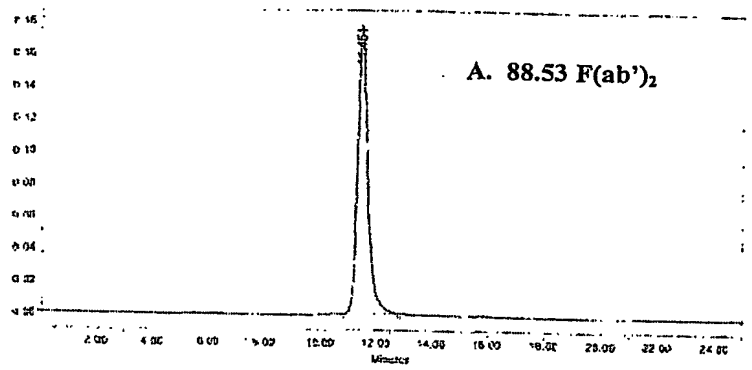


Fig. 7b

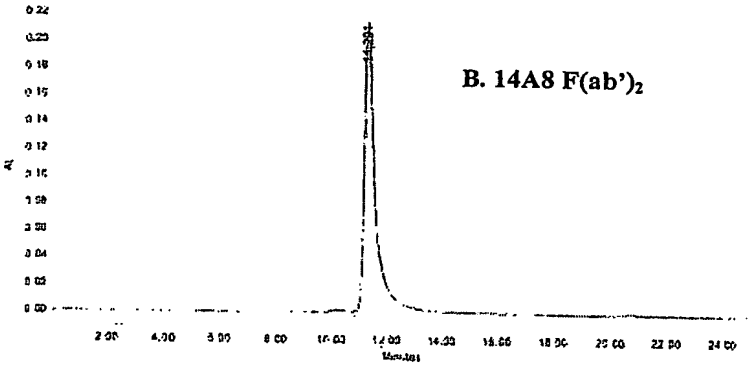


Fig. 8

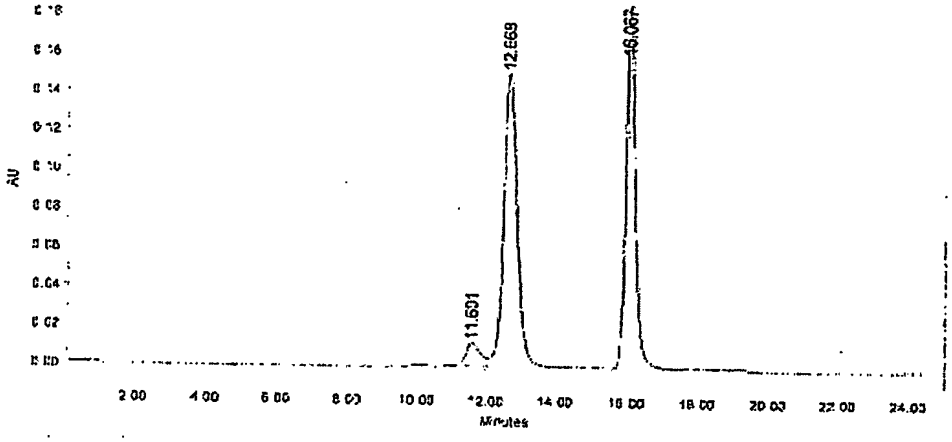


Fig. 9

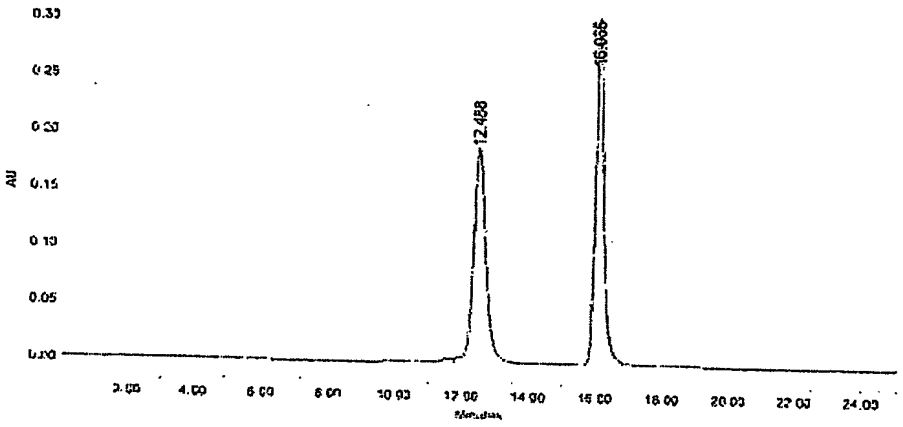


Fig. 10

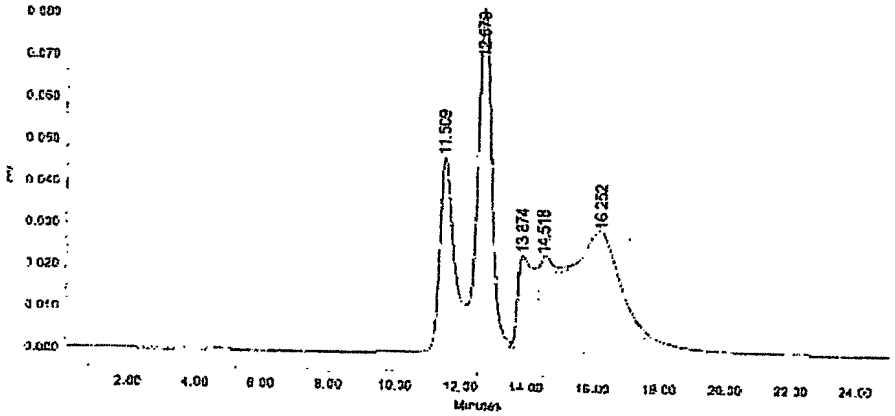


Fig. 11

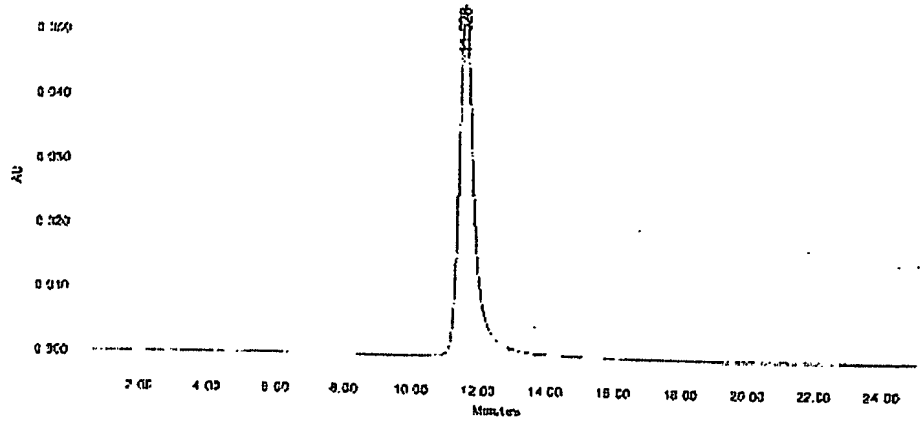


Fig. 12

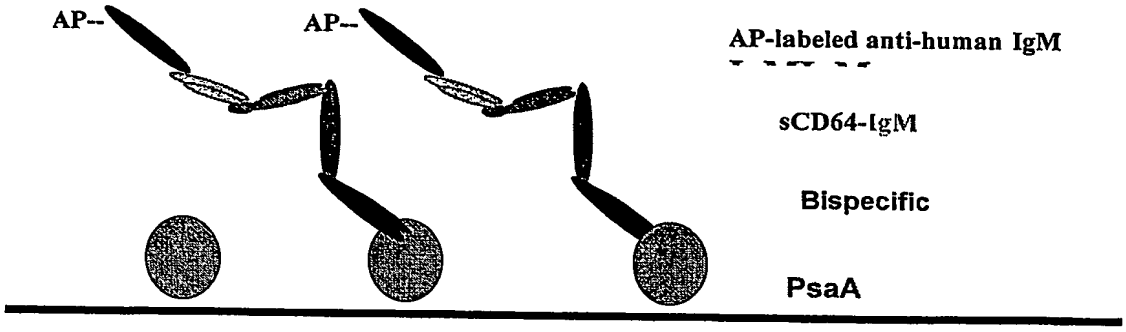


Fig. 13

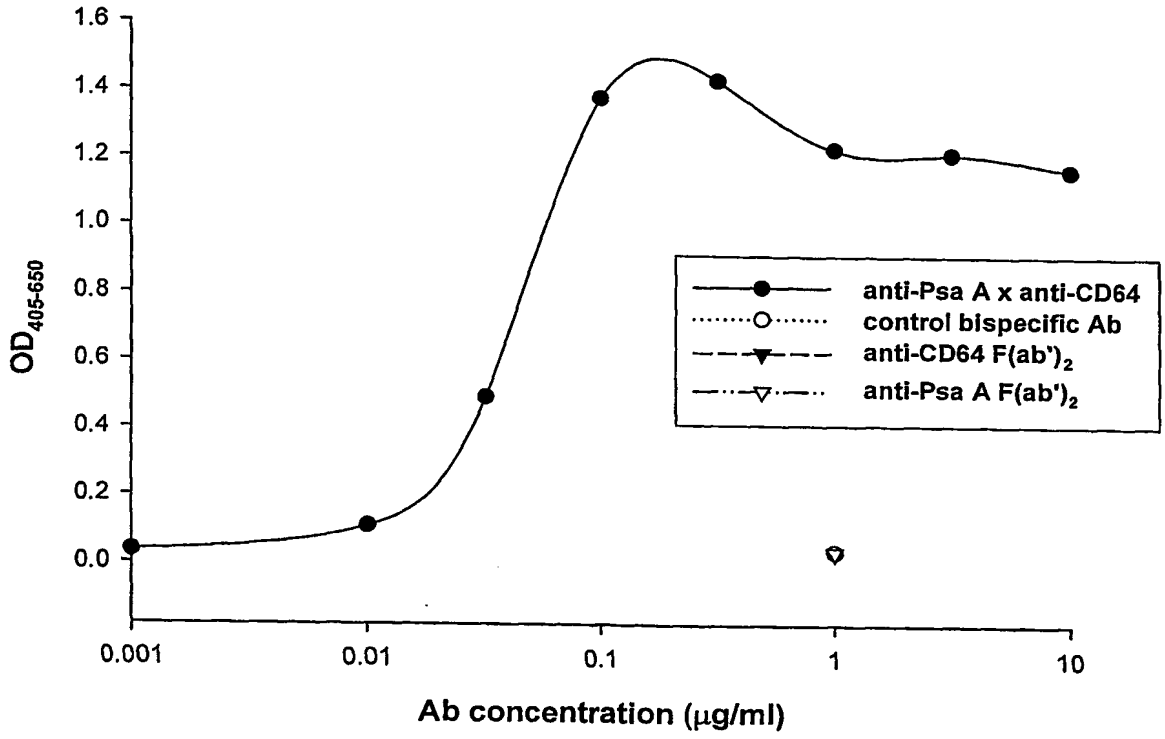

1.6
1.4
1.2
1.0
0.8
0.6
0.4
0.2
0.0
OD405-650
0.001
0.01
0.1
1
10
Ab concentration (µg/ml)
anti-Psa A x anti-CD64
control bispecific Ab
anti-CD64 F(ab')2
anti-Psa A F(ab')2

Fig. 14

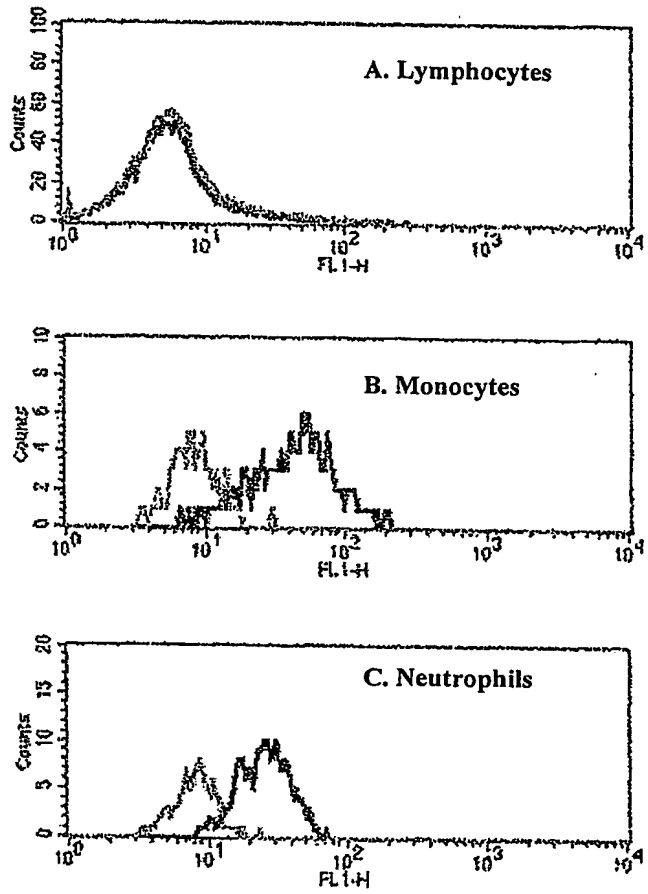

A. Lymphocytes
Counts
FL1-H
B. Monocytes
Counts
FL1-H
C. Neutrophils
Counts
FL1-H

Fig. 15

```
1     ATG AAA AAA TTA GGT ACA TTA CTC GTT CTC TTT CTT TCT GCA ATC   45
1     Met Lys Lys Leu Gly Thr Leu Leu Val Leu Phe Leu Ser Ala Ile   15

46    ATT CTT GTA GCA TGT GCT AGC GGA AAA AAA GAT ACA ACT TCT GGT   90
16    Ile Leu Val Ala Cys Ala Ser Gly Lys Lys Asp Thr Thr Ser Gly   30

91    CAA AAA CTA AAA GTT GTT GCT ACA AAC TCA ATC ATC GCT GAT ATT   135
31    Gln Lys Leu Lys Val Val Ala Thr Asn Ser Ile Ile Ala Asp Ile   45

136   ACT AAA AAT ATT GCT GGT GAC AAA ATT GAC CTT CAT AGT ATC GTT   180
46    Thr Lys Asn Ile Ala Gly Asp Lys Ile Asp Leu His Ser Ile Val   60

181   CCG ATT GGG CAA GAC CCA CAC GAA TAC GAA CCA CTT CCT GAA GAC   225
61    Pro Ile Gly Gln Asp Pro His Glu Tyr Glu Pro Leu Pro Glu Asp   75

226   GTT AAG AAA ACT TCT GAG GCT GAT TTG ATT TTC TAT AAC GGT ATC   270
76    Val Lys Lys Thr Ser Glu Ala Asp Leu Ile Phe Tyr Asn Gly Ile   90

271   AAC CTT GAA ACA GGT GGC AAT GCT TGG TTT ACA AAA TTG GTA GAA   315
91    Asn Leu Glu Thr Gly Gly Asn Ala Trp Phe Thr Lys Leu Val Glu   105

316   AAT GCC AAG AAA ACT GAA AAC AAA GAC TAC TTC GCA GTC AGC GAC   360
106   Asn Ala Lys Lys Thr Glu Asn Lys Asp Tyr Phe Ala Val Ser Asp   120

361   GGC GTT GAT GTT ATC TAC CTT GAA GGT CAA AAT GAA AAA GGA AAA   405
121   Gly Val Asp Val Ile Tyr Leu Glu Gly Gln Asn Glu Lys Gly Lys   135

406   GAA GAC CCA CAC GCT TGG CTT AAC CTT GAA AAC GGT ATT ATT TTT   450
136   Glu Asp Pro His Ala Trp Leu Asn Leu Glu Asn Gly Ile Ile Phe   150

451   GCT AAA AAT ATC GCC AAA CAA TTG AGC GCC AAA GAC CCT AAC AAT   495
151   Ala Lys Asn Ile Ala Lys Gln Leu Ser Ala Lys Asp Pro Asn Asn   165

496   AAA GAA TTC TAT GAA AAA AAT CTC AAA GAA TAT ACT GAT AAG TTA   540
166   Lys Glu Phe Tyr Glu Lys Asn Leu Lys Glu Tyr Thr Asp Lys Leu   180

541   GAC AAA CTT GAT AAA GAA AGT AAG GAT AAA TTT AAT AAG ATC CCT   585
181   Asp Lys Leu Asp Lys Glu Ser Lys Asp Lys Phe Asn Lys Ile Pro   195

586   GCT GAA AAG AAA CTC ATT GTA ACC AGC GAA GGA GCA TTC AAA TAC   630
196   Ala Glu Lys Lys Leu Ile Val Thr Ser Glu Gly Ala Phe Lys Tyr   210

631   TTC TCT AAA GCC TAT GGT GTT CCA AGT GCC TAC ATC TGG GAA ATC   675
211   Phe Ser Lys Ala Tyr Gly Val Pro Ser Ala Tyr Ile Trp Glu Ile   225

676   AAT ACT GAA GAA GAA GGA ACT CCT GAA CAA ATC AAG ACC TTG GTT   720
226   Asn Thr Glu Glu Glu Gly Thr Pro Glu Gln Ile Lys Thr Leu Val   240

721   GAA AAA CTT CGC CAA ACA AAA GTT CCA TCA CTC TTT GTA GAA TCA   765
241   Glu Lys Leu Arg Gln Thr Lys Val Pro Ser Leu Phe Val Glu Ser   255

766   AGT GTG GAT GAC CGT CCA ATG AAA ACT GTT TCT CAA GAC ACA AAC   810
256   Ser Val Asp Asp Arg Pro Met Lys Thr Val Ser Gln Asp Thr Asn   270

811   ATC CCA ATC TAC GCA CAA ATC TTT ACT GAC TCT ATC GCA GAA CAA   855
271   Ile Pro Ile Tyr Ala Gln Ile Phe Thr Asp Ser Ile Ala Glu Gln   285

856   GGT AAA GAA GGC GAC AGC TAC TAC AGC ATG ATG AAA TAC AAC CTT   900
286   Gly Lys Glu Gly Asp Ser Tyr Tyr Ser Met Met Lys Tyr Asn Leu   300

901   GAC AAG ATT GCT GAA GGA TTG GCA AAA TAA   930
301   Asp Lys Ile Ala Glu Gly Leu Ala Lys End
```

Fig. 16a

```
      D   I   Q   M   T   Q   S   P   S   S   L   S   A   S   V   G   D   R
1     GAC ATC CAG ATG ACC CAG TCT CCA TCC TCA CTG TCT GCA TCT GTA GGA GAC AGA

                                CDR1
                            ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
      V   T   I   T   C   R   A   S   Q   G   I   S   S   W   L   A   W   Y
55    GTC ACC ATC ACT TGT CGG GCG AGT CAG GGT ATT AGC AGC TGG TTA GCC TGG TAT

                                                        CDR2
                                                    ~~~~~~~~~~~~~~~~~~~
      Q   Q   K   P   E   K   A   P   E   S   L   I   Y   V   A   S   S   L
109   CAG CAG AAA CCA GAG AAA GCC CCT GAG TCC CTG ATC TAT GTT GCA TCC AGT TTG

      CDR2
      ~~~~~~~~
      Q   S   G   V   P   S   R   F   S   G   S   G   S   G   T   D   F   T
163   CAA AGT GGG GTC CCA TCA AGG TTC AGC GGC AGT GGA TCT GGG ACA GAT TTC ACT

                                                                    CDR3
                                                                    ~~~~~~
      L   T   I   S   S   L   Q   P   E   D   F   A   T   Y   Y   C   Q   Q
217   CTC ACC ATC AGC AGC CTG CAG CCT GAA GAT TTT GCA ACT TAT TAC TGC CAA CAG

              CDR3
      ~~~~~~~~~~~~~~~~~~~~~~~~~~~
      Y   N   S   Y   P   P   T   F   G   Q   G   T   K   V   E   I   K
271   TAT AAT AGC TAT CCT CCG ACG TTC GGC CAA GGG ACC AAG GTG GAA ATC AAA
                    └▷ JK1
```

Fig. 16b

```
      Q   V   R   L   Q   Q   W   G   A   G   L   L   K   P   S   E   T   L
1     CAG GTG CGA CTA CAG CAG TGG GGC GCA GGA CTG TTG AAG CCT TCG GAG ACC CTG

                                                    CDR1
                                                ~~~~~~~~~~~~~~~~~~~~~
      S   L   T   C   A   V   F   G   G   S   F   S   G   F   S   W   S   W
55    TCC CTC ACC TGC GCT GTC TTT GGT GGG TCC TTC AGT GGT TTC TCC TGG AGC TGG

                                                        CDR2
                                                    ~~~~~~~~~~~~~~~~~~~~
      I   R   Q   T   P   G   K   G   L   E   W   I   G   E   I   D   Y   R
109   ATC CGC CAG ACC CCA GGG AAG GGG CTG GAG TGG ATC GGG GAA ATC GAT TAT AGA

                        CDR2
      ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
      G   S   T   N   Y   N   P   S   L   K   S   R   V   T   I   L   R   D
163   GGA AGC ACC AAC TAC AAC CCG TCC CTC AAG AGT CGA GTC ACC ATA TTA AGA GAC

      T   S   R   S   Q   F   S   L   K   L   S   S   V   T   A   A   D   S
217   ACG TCC AGG AGC CAG TTC TCC CTG AAG TTG AGC TCC GTG ACC GCC GCG GAC TCG
```

```
                                                    ~~~~~~~~~~~~~~~~~~~~~~~~~~~~
          A   V   F   Y   C   A   R   G   G   P   R   F   D   Y   W   G   Q   G
271       GCT GTG TTT TAT TGT GCG AGA GGG GGG CCC CGC TTT GAC TAC TGG GGC CAG GGA
                                                    └▷ JH4b

          T   L   V   T   V   S   S
325       ACC CTG GTC ACC GTC TCC TCA
```

Fig. 17a

```
          E   I   V   L   T   Q   S   P   A   T   L   S   L   S   P   G   E   R
1         GAA ATT GTG TTG ACA CAG TCT CCA GCC ACC CTG TCT TTG TCT CCA GGG GAA AGA

                                  CDR1
                              ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
          A   T   L   S   C   R   A   S   Q   S   V   S   S   Y   L   A   W   Y
55        GCC ACC CTC TCC TGC AGG GCC AGT CAG AGT GTT AGC AGC TAC TTA GCN TGG TAC

                                                                  CDR2
                                                              ~~~~~~~~~~~~~~~~~~~
          Q   Q   K   P   G   Q   A   P   R   L   L   I   Y   D   A   S   N   R
109       CAA CAG AAA CCT GGC CAG GCT CCC AGG CTC CTC ATC TAT GAT GCA TCC AAC AGG

              CDR2
          ~~~~~~~~
          A   T   G   I   P   A   R   F   S   G   S   G   S   G   T   D   F   T
163       GCC ACT GGC ATC CCA GCC AGG TTC AGT GGC AGT GGG TCT GGG ACA GAC TTC ACT

                                                                          CDR3
                                                                      ~~~~~~~~
          L   T   I   S   S   L   E   P   E   D   F   A   V   Y   Y   C   Q   Q
217       CTC ACC ATC AGC AGC CTA GAG CCT GAA GAT TTT GCA GTT TAT TAC TGT CAG CAG

                      CDR3
          ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
          R   S   N   W   P   L   T   F   G   G   G   T   K   V   E   I   K
271       CGT AGC AAC TGG CCT CTC ACT TTC GGC GGA GGG ACC AAG GTG GAG ATC AAA
```

Fig. 17b

```
          E   V   Q   L   V   E   S   G   G   G   L   V   Q   P   G   G   S   L
1         GAG GTG CAA CTA GTG GAG TCT GGG GGA GGC TTG GTC CAG CCT GGG GGG TCC CTG

                                                      CDR1
                                                  ~~~~~~~~~~~~~~~~~~~~
          R   L   S   C   A   A   S   G   F   T   F   N   I   F   G   M   S   W
55        AGA CTC TCC TGT GCA GCC TCT GGA TTC ACC TTT AAT ATC TTT GGG ATG AGC TGG

                                                                  CDR2
                                                              ~~~~~~~~~~~~~~~~~~~
          V   R   Q   A   P   G   K   G   L   E   W   V   A   N   I   K   Q   D
109       GTC CGC CAG GCT CCA GGG AAA GGG CTG GAG TGG GTG GCC AAC ATA AAG CAA GAT
```

```
                    CDR2
                    ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
                     G   S   E   K   Y   Y   V   D   S   V   K   G   R   F   T   I   S   R
163                  GGA AGT GAG AAA TAC TAT GTG GAC TCT GTG AAG GGC CGA TTC ACC ATC TCC AGA

                     D   N   A   K   N   S   L   Y   L   Q   M   N   S   L   R   A   E   D
217                  GAC AAC GCC AAG AAC TCA CTG TAT CTG CAA ATG AAC AGC CTG AGA GCC GAG GAC

                                                    CDR3
                                                    ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
                     T   A   V   Y   Y   C   A   R   D   R   F   Y   Y   G   S   G   S   Y
271                  ACG GCT GTG TAT TAC TGT GCG AGG GAT CGG TTT TAC TAT GGT TCG GGG AGT TAT
                                                                                            └▷ JH6b

                             CDR3
                             ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
                     Y   Y   Y   Y   N   G   M   D   V   W   G   Q   G   T   T   V   T   V
325                  TAT TAC TAC TAC AAC GGT ATG GAC GTC TGG GGC CAA GGG ACC ACG GTC ACC GTC

                     S   S
379                  TCC TCA
```

Fig. 18a

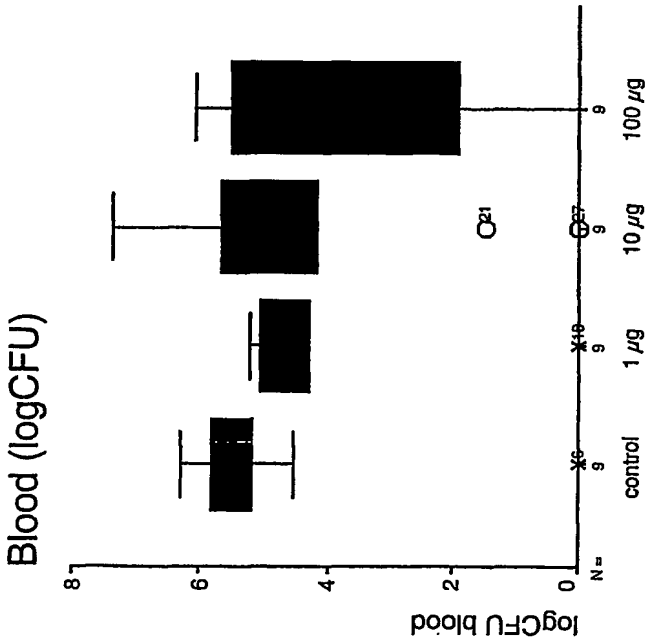
```
                     E   I   V   L   T   Q   S   P   A   T   L   S   L   S   P   G   E   R
1                    GAA ATT GTG TTG ACA CAG TCT CCA GCC ACC CTG TCT TTG TCT CCA GGG GAA AGA

                                    CDR1
                                    ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
                     A   T   L   S   C   R   A   S   Q   S   V   S   S   Y   L   A   W   Y
55                   GCC ACC CTC TCC TGC AGG GCC AGT CAG AGT GTT AGC AGC TAC TTA GCC TGG TAC

                                                                    CDR2
                                                                    ~~~~~~~~~~~~~~~~~~~~
                     Q   Q   K   P   G   Q   A   P   R   L   L   I   Y   D   A   S   N   R
109                  CAA CAG AAA CCT GGC CAG GCT CCC AGG CTC CTC ATC TAT GAT GCA TCC AAC AGG

                     CDR2
                     ~~~~~~~~
                     A   T   G   I   P   A   R   F   S   G   S   G   S   G   T   D   F   T
163                  GCC ACT GGC ATC CCA GCC AGG TTC AGT GGC AGT GGG TCT GGG ACA GAC TTC ACT

                                                                                    CDR3
                                                                                    ~~~~~~~~
                     L   T   I   S   S   L   E   P   E   D   F   A   V   Y   Y   C   Q   Q
217                  CTC ACC ATC AGC AGC CTA GAG CCT GAA GAT TTT GCA GTT TAT TAC TGT CAG CAG

                             CDR3
                             ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
                     R   S   N   W   P   P   F   T   F   G   P   G   T   K   V   D   I   K
271                  CGT AGC AAC TGG CCT CCA TTC ACT TTC GGC CCT GGG ACC AAA GTG GAT ATC AAA
                                                            └▷ JK3
```

Fig. 18b

```
      E   V   Q   L   V   E   S   G   G   G   L   V   Q   P   G   G   S   L
1     GAG GTA CAG CTG GTG GAG TCT GGG GGA GGC TTG GTC CAG CCG GGG GGG TCC CTG

                                                    CDR1
                                                    ~~~~~~~~~~~~~~~~~~~~
      R   L   S   C   A   A   S   G   F   T   F   S   S   F   W   M   S   W
55    AGA CTC TCC TGT GCA GCT TCT GGA TTC ACC TTT AGT AGC TTT TGG ATG AGC TGG

                                                        CDR2
                                                        ~~~~~~~~~~~~~~~~~~~
      V   R   Q   A   P   G   K   G   L   E   W   V   A   N   I   K   Q   D
109   GTC CGC CAG GCT CCA GGG AAG GGG CTG GAG TGG GTG GCC AAC ATA AAG CAA GAT

          CDR2
          ~~~~~~~~~~~~~~~~~~~~
      G   S   E   K   F   Y   V   D   S   V   K   G   R   F   T   I   S   R
163   GGA AGT GAG AAA TTC TAT GTG GAC TCT GTG AAG GGC CGA TTC ACC ATC TCC AGA

      D   N   A   K   N   S   L   Y   L   Q   M   N   S   L   R   A   E   D
217   GAC AAC GCC AAG AAC TCA CTG TAT CTG CAA ATG AAC AGC CTG AGA GCC GAG GAC

                                                CDR3
                                                ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
      T   A   V   Y   Y   C   A   R   D   R   I   T   M   V   R   P   Y   Y
271   ACG GCT GTG TAT TAC TGT GCG AGG GAT CGT ATT ACA ATG GTT CGG CCC TAT TAC
                                                                        └▷

                  CDR3
                                JH6b
      ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
      Y   F   Y   N   G   L   D   V   W   G   Q   G   T   T   V   T   V   S
325   TAC TTC TAC AAC GGT CTG GAC GTC TGG GGC CAA GGG ACC ACG GTC ACC GTC TCC

      S
379   TCA
```

Figure 19A

9A7

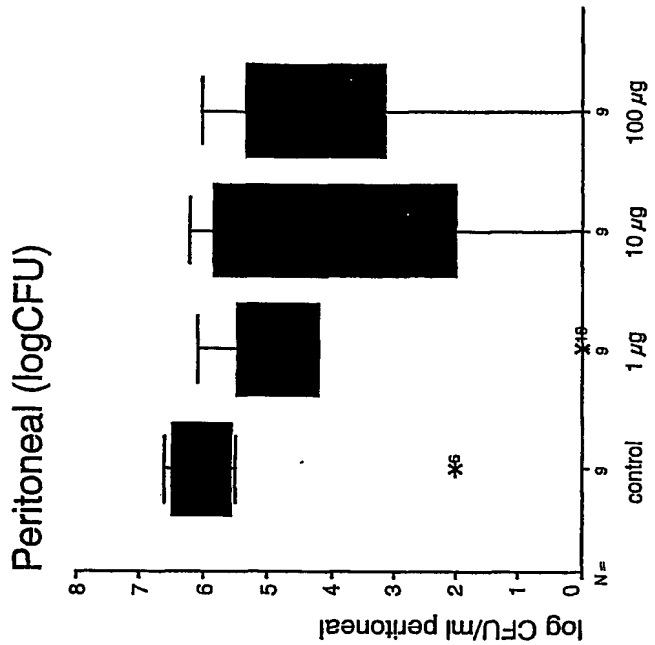


Test Statistics[a,b]

	CFU/ml blood
Chi-Square	4,711
df	3
Asymp. Sig.	,194

a. Kruskal Wallis Test

b. Grouping Variable: AGRP



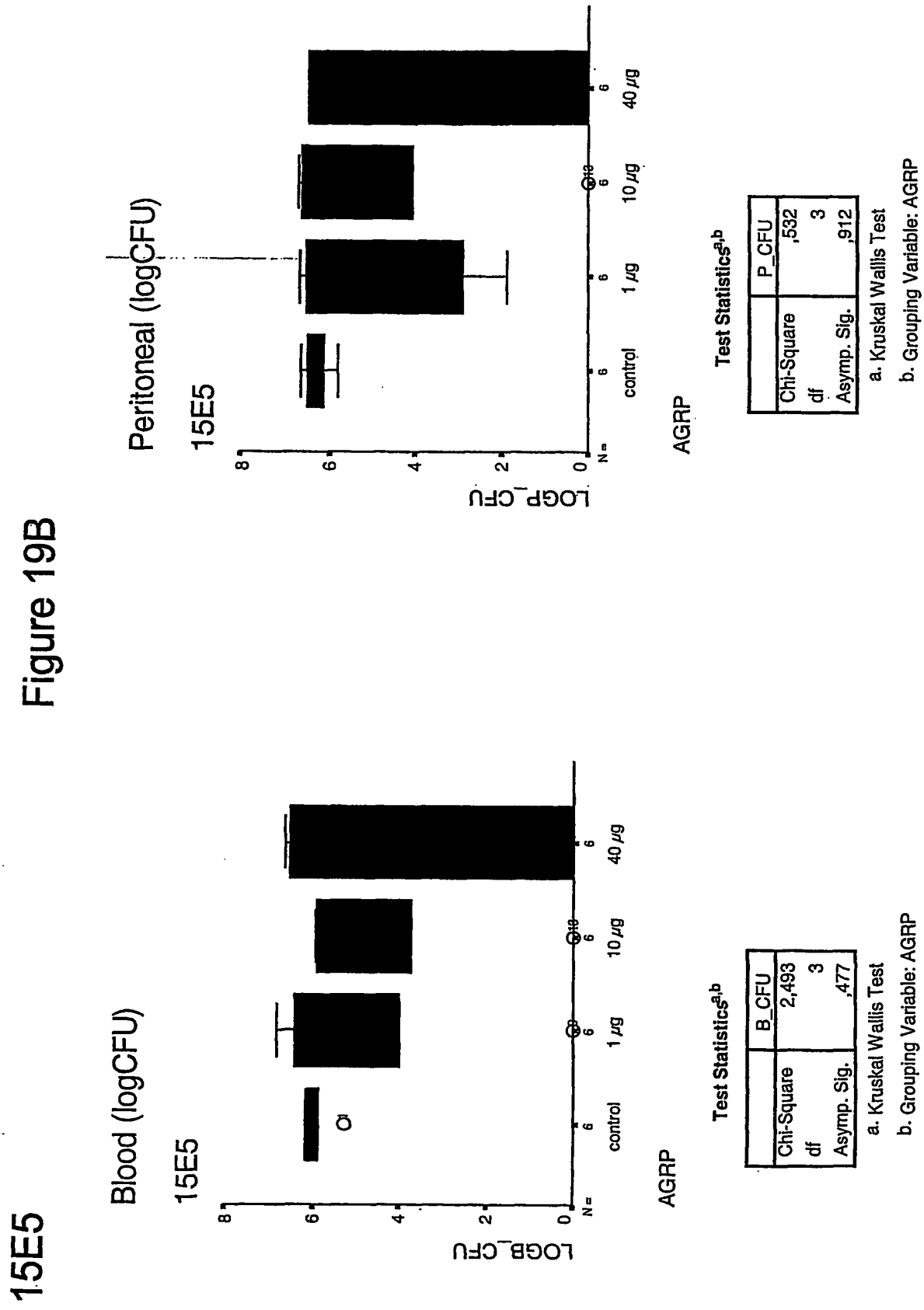
Test Statistics[a,b]

	CFU/ml peritoneal
Chi-Square	7,926
df	3
Asymp. Sig.	,048

a. Kruskal Wallis Test

b. Grouping Variable: AGRP

Figure 19B



Test Statistics[a,b]

	B_CFU
Chi-Square	2,493
df	3
Asymp. Sig.	,477

a. Kruskal Wallis Test

b. Grouping Variable: AGRP

Test Statistics[a,b]

	P_CFU
Chi-Square	,532
df	3
Asymp. Sig.	,912

a. Kruskal Wallis Test

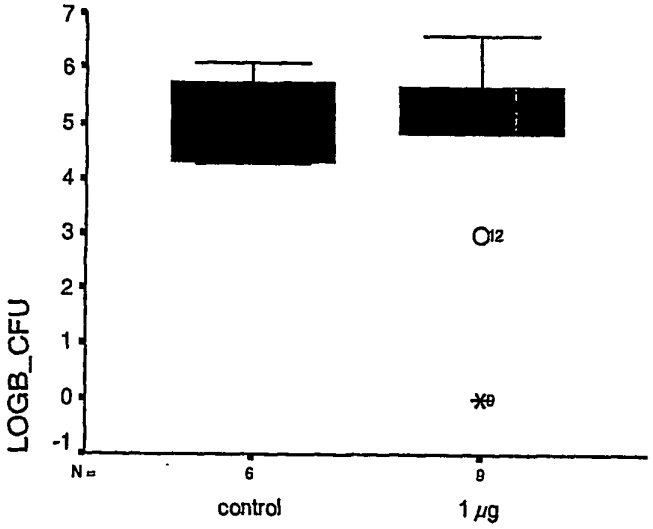
b. Grouping Variable: AGRP

1G9

Figure 19C

Blood (log CFU)

1G9.1



Ab

Test Statistics[b]

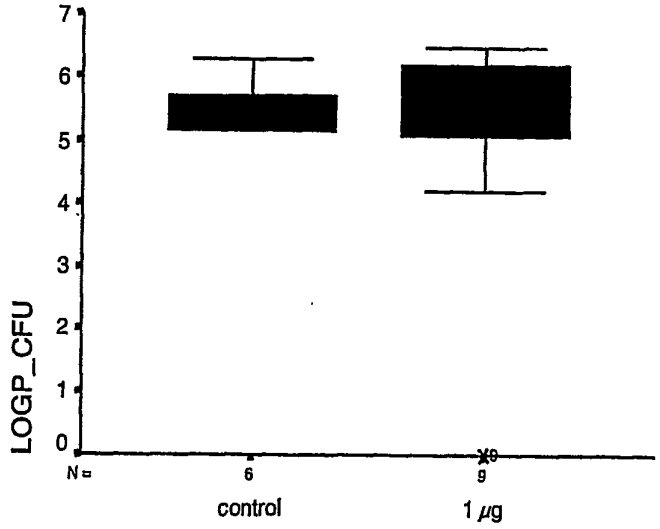
	B_CFU
Mann-Whitney U	23,000
Wilcoxon W	44,000
Z	-,471
Asymp. Sig. (2-tailed)	,637
Exact Sig. [2*(1-tailed Sig.)]	,689[a]

a. Not corrected for ties.

b. Grouping Variable: AGRP

Peritoneal (log CFU)

1G9.1



Ab

Test Statistics[b]

	P_CFU
Mann-Whitney U	22,500
Wilcoxon W	43,500
Z	-,531
Asymp. Sig. (2-tailed)	,596
Exact Sig. [2*(1-tailed Sig.)]	,607[a]

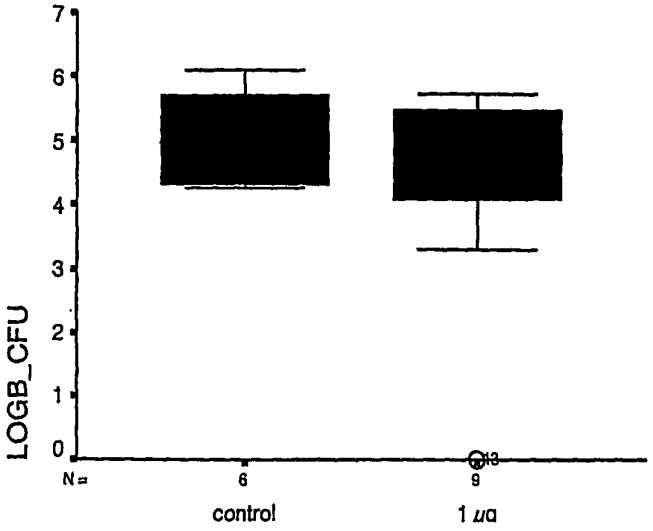
a. Not corrected for ties.

b. Grouping Variable: AGRP

3D10

Figure 19D

Blood (log CFU)



Ab

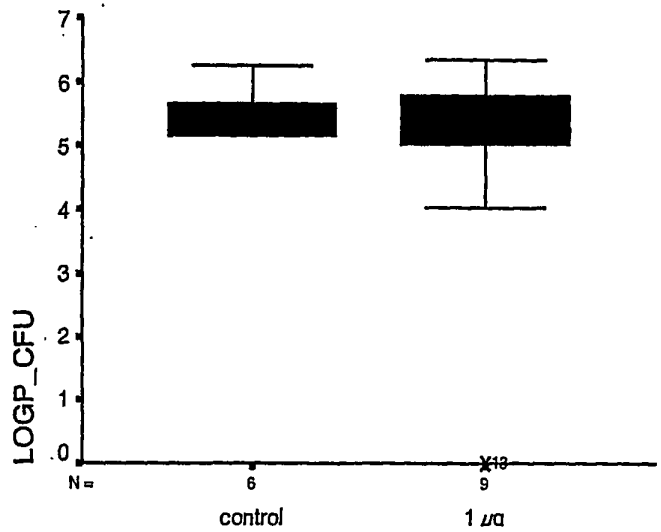
Test Statistics[b]

	B_CFU
Mann-Whitney U	23,000
Wilcoxon W	68,000
Z	-,471
Asymp. Sig. (2-tailed)	,637
Exact Sig. [2*(1-tailed Sig.)]	,689[a]

a. Not corrected for ties.

b. Grouping Variable: AGRP

Peritoneal (log CFU)



Ab

Test Statistics[b]

	P_CFU
Mann-Whitney U	27,000
Wilcoxon W	72,000
Z	,000
Asymp. Sig. (2-tailed)	1,000
Exact Sig. [2*(1-tailed Sig.)]	1,000[a]

a. Not corrected for ties.

b. Grouping Variable: AGRP

Figure 19E

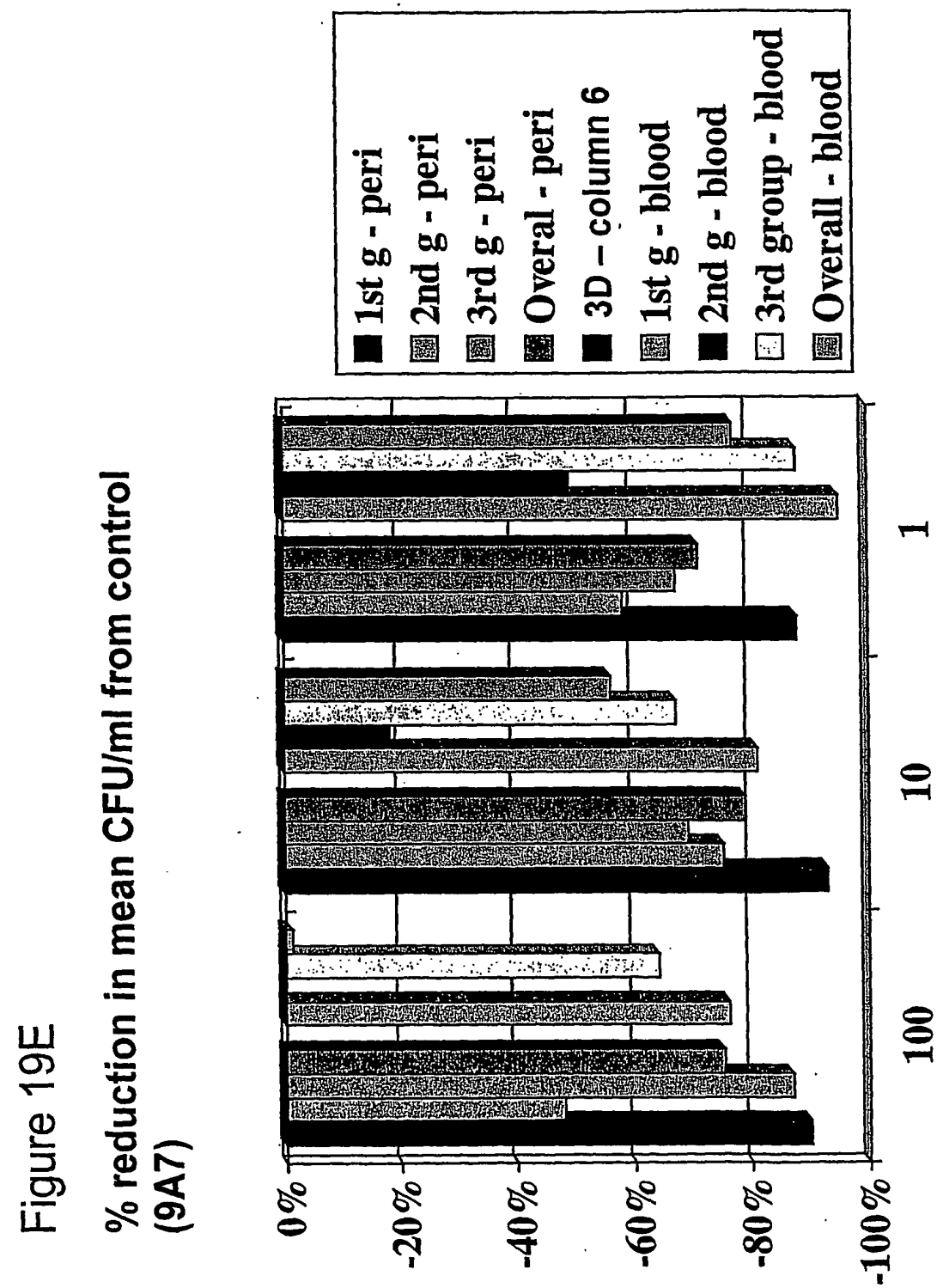

% reduction in mean CFU/ml from control (9A7)
0%
-20%
-40%
-60%
-80%
-100%
1
10
100
1st g - peri
2nd g - peri
3rd g - peri
Overal - peri
3D – column 6
1st g - blood
2nd g - blood
3rd group - blood
Overall - blood

Fig. 20

Clone	Sub type	c1q	R6 ELISA (corrected)					R6 WB	ELISA (0.5 µg/ml) (corrected)												Western Blot (2 µg/ml)								
			5 µg/ml	0.5 µg/ml	50 ng/ml	5 ng/ml	0.5 ng/ml	2 µg/ml	R6	1	3	4	6B	7F	8	9V	12F	14	23F	D39	R6	1	3	4	6B	7F	8	9V	12F
1-7D12	G3		1,58	0,35	0,23	0,20	0,20	0	0,42	0,45	0,38	0,40	0,60	0,49	0,55	0,44	0,51	0,38	0,40	1,23	0	0	0	0	0	0	0	0	0
1-7H7	G3		0,00	0,50	0,22	0,19	0,20	0	0,38	0,52	0,40	0,49	0,64	0,53	0,62	0,50	0,60	0,39	0,44	0,69	0	0	0	0	0	0	0	0	0
1-11H10	G3		0,00	0,86	0,52	0,26	0,22	1	0,90	0,30	0,79	0,26	0,39	0,81	1,08	2,56	1,10	0,77	0,24	0,06	1	1	1	1	1	1	1	1	1
1-12E10	G3		2,01	0,42	0,24	0,21	0,21	0	0,56	0,47	0,29	0,31	0,48	0,51	0,46	0,55	0,50	0,38	0,37	0,83	0	0	0	0	0	0	0	0	0
1-15E5	G1		1,74	1,71	0,79	0,30	0,25	1	0,99	0,31	0,97	0,30	0,37	0,96	1,19	2,49	1,09	1,09	0,33	0,68	1	1	1	1	1	1	1	1	1
1-15E5.2.1	**G1**		**1,33**	**1,69**	**1,53**	**0,55**	**0,32**	**1**	**1,20**	**0,36**	**1,06**	**0,35**	**0,35**	**1,06**	**1,33**	**>3**	**1,23**	**1,14**	**0,34**	**0,65**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**
3-4F10	G1		0,98	0,38	0,24	0,21	0,17	1	0,45	0,53	0,37	0,60	0,59	0,46	0,42	0,75	0,38	0,66	0,41	0,72	1	1	1	1	(0)	1	1	1	1
4-3D10	**G3**		**1,23**	**1,19**	**0,79**	**0,33**	**0,25**	**1**	**0,90**	**0,33**	**0,92**	**0,33**	**0,44**	**0,74**	**0,95**	**2,31**	**1,06**	**0,92**	**0,37**	**0,91**	**1**	**1**	**1**	**1**	**(0)**	**1**	**1**	**1**	**1**
4-4B11	G3		1,23	0,90	0,49	0,29	0,26	1	0,83	0,36	0,79	0,34	0,37	0,66	0,87	1,93	0,86	0,67	0,29	0,81	1	1	1	1	(0)	1	1	1	1
4-7E8	G3		0,00	-0,45	0,29	0,27	0,28	1	-1,46	-2,00	-1,46	-1,96	-1,94	-1,55	-1,28	-0,23	-1,23	-1,40	-1,97	-1,23	1	1	1	1	(0)	1	1	1	1
4-7H8	G3		1,45	1,29	0,54	0,31	0,27	1	0,74	0,26	0,66	0,20	0,31	0,63	0,84	2,04	0,87	0,66	0,28	0,58	1	1	1	1	(0)	1	1	1	1
4-10G6	G1		0,97	0,35	0,25	0,24	0,25	0	0,32	0,07	0,24	0,07	0,19	0,29	0,35	0,36	0,36	0,25	0,10	0,41	1	1	1	1	(0)	1	1	1	1
5-2G6	G1		0,58	0,42	0,31	0,25	0,25	1	0,47	0,07	0,27	0,06	0,18	0,29	0,40	1,05	0,38	0,20	0,09	0,38	1	1	1	1	(0)	1	1	1	1
5-2G8	**G1**		**1,57**	**1,61**	**0,80**	**0,33**	**0,27**	**1**	**1,38**	**0,35**	**1,18**	**0,28**	**0,34**	**0,93**	**1,24**	**2,83**	**1,27**	**1,05**	**0,30**	**0,62**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**
5-5E10	G1		1,74	1,59	0,57	0,29	0,26	1	1,33	0,31	1,28	0,30	0,34	0,93	1,33	2,74	1,24	1,03	0,31	0,59	1	1	1	1	1	1	1	1	1
5-9A7	**G1**		**1,39**	**1,65**	**1,32**	**0,41**	**0,26**	**1**	**1,39**	**0,40**	**1,31**	**0,33**	**0,39**	**1,17**	**1,52**	**>3**	**1,51**	**1,42**	**0,37**	**0,66**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**
5-9C7	G1		1,65	0,88	0,35	0,26	0,24	1	1,14	0,26	0,86	0,20	0,33	0,82	1,12	2,29	1,10	0,95	0,27	0,65	1	1	1	1	1	1	1	1	1
5-10G9	G1		0,97	0,77	0,36	0,26	0,26	1	0,25	0,03	0,02	0,03	0,16	0,18	0,19	0,13	0,17	0,05	0,06	0,40	1	1	1	1	1	1	1	1	1
6-7A4	**G1**		**1,38**	**1,69**	**1,08**	**0,43**	**0,28**	**1**	**1,20**	**0,24**	**0,96**	**0,19**	**0,29**	**0,71**	**1,11**	**2,66**	**1,04**	**0,85**	**0,25**	**0,63**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**
6-10E5	G1		-0,11	1,12	0,86	0,33	0,22	1	1,32	0,35	1,14	0,32	0,39	1,05	1,44	>3	1,44	1,41	0,38	0,52	1	1	1	1	1	1	1	1	1
7-1G9.1	**G1**		**1,36**	**1,68**	**1,78**	**0,54**	**0,31**	**1**	**1,38**	**0,35**	**1,24**	**0,33**	**0,40**	**1,06**	**1,50**	**>3**	**1,46**	**1,32**	**0,36**	**0,68**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**	**1**
7-7F12.1	G1		0,00	1,01	1,08	0,39	0,27	1	1,06	0,25	0,99	0,27	0,37	0,94	1,15	>3	1,27	1,06	0,31	0,39	1	1	1	1	1	1	1	1	1
7-9E2.1	G1		0,00	1,10	1,18	0,47	0,29	1	1,01	0,28	0,91	0,24	0,29	0,93	1,13	2,46	1,30	1,05	0,29	0,45	1	1	1	1	1	1	1	1	1